



PCT09

RAW SEQUENCE LISTING

DATE: 01/22/2003

PATENT APPLICATION: US/09/647,481A

TIME: 14:55:36

Input Set : A:\Ahmad.ST25.txt

Output Set: N:\CRF4\01222003\I647481A.raw

3 <110> APPLICANT: Ahmad, Sultan
 4 Hoffert, Cyria
 5 O'Donnell, Dajan
 6 Pelletier, Manon
 7 Walker, Philippe
 9 <110> TITLE OF INVENTION: Receptor
 11 <130> FILE REFERENCE: 7507/73170
 13 <140> CURRENT APPLICATION NUMBER: 09/647,481A
 14 <141> CURRENT FILING DATE: 2000-09-29
 16 <160> NUMBER OF SEQ ID NOS: 6
 18 <170> SOFTWARE: PatentIn version 3.1
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 400
 22 <212> TYPE: PRT
 23 <213> ORGANISM: Rattus norvegicus
 25 <400> SEQUENCE: 1
 27 Met Glu Ser Gly Leu Leu Arg Pro Ala Pro Val Ser Glu Val Ile Val
 28 1 5 10 15
 31 Leu His Tyr Asn Tyr Thr Gly Lys Leu Arg Gly Ala Arg Tyr Gln Pro
 32 10 25 30
 35 Gly Ala Gly Leu Arg Ala Asp Ala Ala Val Cys Leu Ala Val Cys Ala
 36 35 40 45
 39 Phe Ile Val Leu Glu Asn Leu Ala Val Leu Leu Val Leu Gly Arg His
 40 50 55 60
 43 Pro Arg Phe His Ala Pro Met Phe Leu Leu Leu Gly Ser Leu Thr Leu
 44 65 70 75 80
 47 Ser Asp Leu Leu Ala Gly Ala Ala Tyr Ala Thr Asn Ile Leu Leu Ser
 48 85 90 95
 51 Gly Pro Leu Thr Leu Arg Leu Ser Pro Ala Leu Trp Phe Ala Arg Glu
 52 100 105 110
 55 Gly Gly Val Phe Val Ala Leu Ala Ala Ser Val Leu Ser Leu Leu Ala
 56 115 120 125
 59 Ile Ala Leu Glu Arg His Leu Thr Met Ala Arg Arg Gly Pro Ala Pro
 60 130 135 140
 63 Ala Ala Ser Arg Ala Arg Thr Leu Ala Met Ala Val Ala Ala Trp Gly
 64 145 150 155 160
 67 Leu Ser Leu Leu Leu Gly Leu Leu Pro Ala Leu Gly Trp Asn Cys Leu
 68 165 170 175
 71 Gly Arg Leu Glu Ala Cys Ser Thr Val Leu Pro Leu Tyr Ala Lys Ala
 72 180 185 190
 75 Tyr Val Leu Phe Cys Val Leu Ala Phe Leu Gly Ile Leu Ala Ala Ile
 76 195 200 205
 79 Cys Ala Leu Tyr Ala Arg Ile Tyr Cys Gln Val Arg Ala Asn Ala Arg

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80      210      215      220
83 Arg Leu Arg Ala Gly Pro Gly Ser Arg Arg Ala Thr Ser Ser Ser Arg
84 325      330      335      340
87 Ser Arg His Thr Pro Arg Ser Leu Ala Leu Leu Arg Thr Leu Ser Val
88      345      250      255
91 Val Leu Leu Ala Phe Val Ala Cys Trp Gly Pro Leu Phe Leu Leu Leu
92      260      265      270
95 Leu Leu Asp Val Ala Cys Pro Ala Arg Ala Cys Pro Val Leu Leu Gln
96      275      280      285
99 Ala Asp Pro Phe Leu Gly Leu Ala Met Ala Asn Ser Leu Leu Asn Pro
100      290      295      300
103 Ile Ile Tyr Thr Phe Thr Asn Arg Asp Leu Arg His Ala Leu Leu Arg
104 305      310      315      320
107 Leu Leu Cys Cys Gly Arg Gly Pro Cys Asn Gln Asp Ser Ser Asn Ser
108      325      330      335
111 Leu Gln Arg Ser Pro Ser Ala Val Gly Pro Ser Gly Gly Gly Leu Arg
112      340      345      350
115 Arg Cys Leu Pro Thr Thr Leu Asp Arg Ser Ser Ser Pro Ser Gln His
116      355      360      365
119 Ser Cys Pro Gln Arg Asp Gly Met Asp Thr Ser Cys Ser Thr Gly Ser
120      370      375      380
123 Pro Gly Ala Ala Thr Ala Asn Arg Thr Leu Val Pro Asp Ala Thr Asp
124 385      390      395      400
127 <C10> SEQ ID NO: 1
128 <C11> LENGTH: 120
129 <C12> TYPE: DNA
130 <C13> ORGANISM: Ectus norvegicus
132 <C14> SEQUENCE:

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133 atggactcgg ggcactcggc ggcagcgcgc gtgagcgggg tcctcgtcct tcaactcaac 60
135 tacactggca agctccggcg aagcgctac cagcccggtg ccggcctcgc tgcggaagcc 120
137 gcaatgtgct tgcctgggag cgtcttcacc gtgctcgaga acctggcgtg gctcttggtg 180
139 ctgggcgcgc atctcgcctt ccactgcacc cgtctcctgc tcttggttag tcttaccctg 240
141 tgggaactgc tctcgcgggc ggcctacgac accaacatcc tgcctcgcgg gccgtcaca 300
143 ttgcgcctgt cgcctgcctt cgtctttgag cgcgaagggg gtgtcttctg ggccctcgca 360
145 ggcctcgggc tgcctcctct ggccttctgt ctgagcgcgc accttaccat ggcccgctgt 420
147 ggaacccgac ctgcacccag tgcgcctcgc accgtggcga tggcggtagc cgcctggggc 480
149 ttgcctcgc tgcctggcct gctggccggg ctgggcctga actgcttggg acccctggaa 540
151 gcctgcctca cgcctcgcgc gctctacgac aaggcctatg tgcctctctg cgtgcgggac 600
153 ttcctgggca tctcgcctgc cctctctgag ctctatgcaa ggatitactg tcaggttgcg 660
155 gcaaacgcgc ctgcgcgggc ggcgggtcct gggtcccgta gggccacgta ctctcccgca 720
157 tcccggcaca ggcacccggt cttgcgcctg ctccgcacgc ttgcgctggt gctcctgggc 780
159 ttgcctggct gctgggaacc tctatctctc ttgcctattc tggatctcgc gtcgcacgac 840
161 cgcgcctgtc ctgctcctct gaaagcgcct cctctcctgg ctctacccat ggctaacctg 900
163 ctgctgaate ctatccatca cacttccacc aacccgaccc tgcgcacgca gctcctgagg 960
165 ctgctctgct gtcgcgcggg accttgcacc caagactcct ccaacagttt gcagcgatcc 1020
167 ccaagtgtcg ttgcaccttc cgttggagag ctgcgagcgt gcttgcaccc aacctggat 1080
169 cgcagctcta ggccttcaga acaactcgtt ccccgagggg accgcatgga caccagctgc 1140
171 tcaactggca gtcgcgggag agcaacgcgc aacccgaccc tggctgcctga tgcctacagc 1200
173 tga

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176 ->110- SEQ ID NO: 3
177 ->111- LENGTH: 12
178 ->112- TYPE: DNA
179 ->113- ORGANISM: Rattus norvegicus
181 ->400- SEQUENCE: 3
182 atttaggtga cactatagaa ta
185 ->110- SEQ ID NO: 4
186 ->111- LENGTH: 19
187 ->112- TYPE: DNA
188 ->113- ORGANISM: Rattus norvegicus
190 ->400- SEQUENCE: 4
191 gctgggggag gactagatg
194 ->110- SEQ ID NO: 5
195 ->111- LENGTH: 20
196 ->112- TYPE: DNA
197 ->113- ORGANISM: Rattus norvegicus
199 ->400- SEQUENCE: 5
200 cctctagatg cagctcgag
203 ->110- SEQ ID NO: 6
204 ->111- LENGTH: 19
205 ->112- TYPE: DNA
206 ->113- ORGANISM: Rattus norvegicus
208 ->400- SEQUENCE: 6
209 caggagcagg ccaaacagg

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VERIFICATION SUMMARY

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